rage 1 of /

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RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/509,945

DATE: 07/30/2001 TIME: 15:59:23

Input Set : A:\0230-0148P.ST25.txt
Output Set: N:\CRF3\07302001\1509945.raw

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3 <110> APPLICANT: HAMADA, Kazuvuki et al.
      5 <120> TITLE OF INVENTION: MUTANT BARNASE GENE AND TRANSGENIC PLANT TRANSFORMED BY SAID
CENE
      7 <130> FILE REFERENCE: 0230-0148P
      9 <140> CURRENT APPLICATION NUMBER: 09/509,945
C--> 10 <141> CURRENT FILING DATE: 2001-04-01
     12 <160> NUMBER OF SEC ID NOS: 7
     14 <170> SOFTWARE: PatentIn version 3.1
                                                               ENTERED
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     29 <223> OTHER INFORMATION:
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                                           10
                                                                              96
     37 aca tat cat aag cta cct gat aat tac att aca aaa tca gaa gca caa
     38 Thr Tyr His Lys Leu Pro Asp Asn Tyr Ile Thr Lys Ser Glu Ala Gln
                    20
                                       25
     41 gcc ctc ggc tgg gtg gca tca aaa ggg aac ctt gca gac gtc gct ccg
                                                                             144
     42 Ala Leu Gly Trp Val Ala Ser Lys Gly Asn Leu Ala Asp Val Ala Pro
               35
                                   40
     43
     45 ggg aaa agc atc ggc gga gac atc ttc tca aac agg gaa ggc aaa ctc
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     46 Gly Lys Ser Ile Gly Gly Asp Ile Phe Ser Asn Arg Glu Gly Lys Leu
                               55
     47
     49 ccg ggc aaa agc gga cga aca tgg cgt gaa gcg gat att aac tat aca
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     50 Pro Gly Lys Ser Gly Arg Thr Trp Arg Glu Ala Asp Ile Asn Tyr Thr
                                               75
     53 tca qqc ttc aga aat tca qac cgg att ctt tac tca agc gac tgg ctg
                                                                             288
     54 Ser Gly Phe Arg Asn Ser Asp Arg Ile Leu Tyr Ser Ser Asp Trp Leu
                                           90
                       85
     57 att tac aaa aca acg gac cat tat cag acc ttt aca aaa atc aga taa
    58 Ile Tyr Lys Thr Thr Asp His Tyr Gln Thr Phe Thr Lys Ile Arg
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    61 ggtaacc
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    65 <211> LENGTH: 111
    66 <212> TYPE: PRT
    67 <213> ORGANISM: Bacillus amyloliquefaciens
    69 <220> FEATURE:
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70 <221> NAME/KEY: misc\_feature

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                                                            30
    RΛ
    83 Ala Leu Gly Trp Val Ala Ser Lys Gly Asn Leu Ala Asp Val Ala Pro
                                                        45
                                    40
               35
     87 Gly Lys Ser Ile Gly Gly Asp Ile Phe Ser Asn Arg Glu Gly Lys Leu
                                55
            50
     91 Pro Gly Lys Ser Gly Arg Thr Trp Arg Glu Ala Asp Ile Asn Tyr Thr
                            70
                                                75
     92 65
     95 Ser Gly Phe Arg Asn Ser Asp Arg Ile Leu Tyr Ser Ser Asp Trp Leu
                                            an
                        85
     99 Ile Tyr Lys Thr Thr Asp His Tyr Gln Thr Phe Thr Lys Ile Arg
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     106 <213> ORGANISM: Artificial Sequence
     108 <220> FEATURE:
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                                                                               120
     116 agggaacett geagacgteg eteeggggaa aageategge ggagacatet teteaaacag
                                                                               180
     118 qqaaqqcaaa ctcccgggca aaaqcggacg aacatggcgt gaagcggata ttaactatac
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     120 atcaggette agaaatteag accggattet ttacteaage gaetggetga tttacaaaac
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     140 cttcaataat attgaaaaag gaagagtatg agtattcaac atttccgtgt cgcccttatt
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     142 cccttttttg cgccattttg ccttcctgtt tttgctcacc cagaaacgct ggtgaaagta
     144 aaagatgctg aagatcagtt gggtgcacga gtgggttaca tcgaactgga tctcaacagc
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     146 ggtaagatcc ttgagagttt tcgccccgaa gaacgttttc caatgatgag cacttttaaa
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     148 gttctgctat gtggcgcggt attatcccgt attgacgccg ggcaaqaqca actcgqtcqc
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     150 cgcatacact attctcagaa tgacttggtt gagtactcac cagtcacaga aaagcatctt
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     152 acggatggca tgacagtaag agaattatgc agtgctgcca taaccatgag tgataacact
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     154 gcggccaact tacttctgac aacgatcgga ggaccgaagg agctaaccgc ttttttgcac
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     156 aacatqqqqq atcatgtaac tcgccttgat cgttgggaac cggagctgaa tgaagccata
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158 ccaaacgacg agcgtgacac cacgatgcct gtagcaatgg caacaacgtt gcgcaaacta

720

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PATENT APPLICATION: US/09/509,945

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					cagcactggg		900
166	aagecetece	gtatcgtagt	tatctacacq	acggggagtc	aggcaactat	ggatgaacga	960
168	aatagacaga	tegetgagat	aggtgcctca	ctgattaagc	attggtaact	gtcagaccaa	1020
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172	gtgaagatcc	tttttggctc	gagtctcatg	accaaaatcc	cttaacgtga	gttttcgttc	1140
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176	cacataatet	actacttaca	aacaaaaaa	ccaccactac	cagcggtggt	ttatttacca	1260
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182	cctacatacc	teactetact	aatcctgtta	ccagtggctg	ctgccagtgg	cgataagtcg	1440
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194	tactcatcaa	aaaaacaaaa	cctatggaaa	aacgccagca	acgcggcctt	tttacggttc	1800
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					gccgcagccg		1920
					tacgcaaacc		1980
202	acacattaac	ctgatcagaa	ttcatatgca	catattccca	atctagtaac	atagatgaca	2040
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246	tcctcatttt	ccgagagatt	ctgacagtga	ccagaatgtc	agaatgccat	ttcatgggca	3360
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266	tctttacgac	attgcatgtg	gaaaggatct	gaagagattt	ctcctggtac	ataataatct	3960
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274	actcaatata	gttctggact	aacaatcaga	ttgtttatga	tattaaggtg	gttggatete	4200
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282	agaatcgggt	attcaacacc	ctcatatcaa	ctactacgtt	gtgtataacg	gtccacatgc	4440
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354 gtatcgcg

6548

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VERIFICATION SUMMARY

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L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date